

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/735,712

DATE: 06/21/2001

TIME: 17:56:57

Input Set : A:\SEQLIST - LEX-0109-USA.txt
Output Set: N:\CRF3\06212001\I735712.raw

ENTERED

4 <110> APPLICANT: Walke, D. Wade
5 Turner, C. Alexander Jr.
7 <120> TITLE OF INVENTION: Novel Human Membrane Proteins and
8 Polynucleotides Encoding the Same
11 <130> FILE REFERENCE: LEX-0109-USA
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/735,712
C--> 13 <141> CURRENT FILING DATE: 2000-12-12
13 <150> PRIOR APPLICATION NUMBER: US 60/171,567
14 <151> PRIOR FILING DATE: 1999-12-22
16 <160> NUMBER OF SEQ ID NOS: 9
18 <170> SOFTWARE: FastSEQ for Windows Version 4.0
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 603
22 <212> TYPE: DNA
23 <213> ORGANISM: Homo sapien
25 <400> SEQUENCE: 1
26 atggattcaa gcaccgcaca cagtcgggtg tttctggat ttcctccaga aatcaactgct 60
27 tcagaatatg agtccacaga actttcagcc acgacacctt caactcaaag ccccttgcaa 120
28 aaattatgg ctagaaaaat gaaaatctta gggactatcc agatcctgtt tggaattatg 180
29 acctttctt ttggagttat cttcctttc acctgttta aaccatatcc aaggtttccc 240
30 tttataatttc ttccagggata tccattctgg ggctctgttt tgttcattaa ttctggagcc 300
31 ttccctaattt cagtggaaaag aaaaaccaca gaaaactctga taatatttgag ccgaataatg 360
32 aatcttctta gtgccttgag agcaatagct ggaatcattc tcctcacatt tggtttcatc 420
33 ctagatcaa actacattt cgttattctt caccaaaata gtcagtgtaa ggctgttact 480
34 gtccctgttct tggaaatttt gattacattt atgactttca gcattatttga attatttattt 540
35 tctctgcctt tctcaattttt ggggtgccac tcagaggatt gtgattgtga acaatgttgt 600
36 tga 603
38 <210> SEQ ID NO: 2
39 <211> LENGTH: 199
40 <212> TYPE: PRT
41 <213> ORGANISM: Homo sapien
43 <400> SEQUENCE: 2
44 Met Asp Ser Ser Thr Ala His Ser Pro Val Phe Leu Val Phe Pro Pro
45 1 5 10 15
46 Glu Ile Thr Ala Ser Glu Tyr Glu Ser Thr Glu Leu Ser Ala Thr Thr
47 20 25 30
48 Phe Ser Thr Gln Ser Pro Leu Gln Lys Leu Phe Ala Arg Lys Met Lys
49 35 40 45
50 Ile Leu Gly Thr Ile Gln Ile Leu Phe Gly Ile Met Thr Phe Ser Phe
51 50 55 60
52 Gly Val Ile Phe Leu Phe Thr Leu Leu Lys Pro Tyr Pro Arg Phe Pro
53 65 70 75 80
54 Phe Ile Phe Leu Ser Gly Tyr Pro Phe Trp Gly Ser Val Leu Phe Ile
55 85 90 95
56 Asn Ser Gly Ala Phe Leu Ile Ala Val Lys Arg Lys Thr Thr Glu Thr
57 100 105 110
58 Leu Ile Ile Leu Ser Arg Ile Met Asn Leu Leu Ser Ala Leu Arg Ala

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59	115	120	125	
60	Ile Ala Gly Ile Ile Leu Leu Thr Phe Gly Phe Ile Leu Asp Gln Asn			
61	130	135	140	
62	Tyr Ile Cys Gly Tyr Ser His Gln Asn Ser Gln Cys Lys Ala Val Thr			
63	145	150	155	160
64	Val Leu Phe Leu Gly Ile Leu Ile Thr Leu Met Thr Phe Ser Ile Ile			
65	165	170	175	
66	Glu Leu Phe Ile Ser Leu Pro Phe Ser Ile Leu Gly Cys His Ser Glu			
67	180	185	190	
68	Asp Cys Asp Cys Glu Gln Cys			
69	195			
71	<210> SEQ ID NO: 3			
72	<211> LENGTH: 246			
73	<212> TYPE: DNA			
74	<213> ORGANISM: Homo sapien			
76	<400> SEQUENCE: 3			
77	atgaatcttc tttagtgcctt gagagcaata gctggaaatca ttctcctcac atttggtttc		60	
78	atccttagatc aaaactacat ttgtggttat ttcacccaaa atagtcagtg taaggctgtt		120	
79	actgtcctgt tcttggaaat tttgattaca ttgatgactt tcagcattat tgaattatttc		180	
80	atttctctgc ctttctcaat tttgggggtgc cactcagagg attgtgattt tgaacaatgt		240	
81	tgttga		246	
83	<210> SEQ ID NO: 4			
84	<211> LENGTH: 81			
85	<212> TYPE: PRT			
86	<213> ORGANISM: Homo sapien			
88	<400> SEQUENCE: 4			
89	Met Asn Leu Leu Ser Ala Leu Arg Ala Ile Ala Gly Ile Ile Leu Leu			
90	1 5 10 15			
91	Thr Phe Gly Phe Ile Leu Asp Gln Asn Tyr Ile Cys Gly Tyr Ser His			
92	20 25 30			
93	Gln Asn Ser Gln Cys Lys Ala Val Thr Val Leu Phe Leu Gly Ile Leu			
94	35 40 45			
95	Ile Thr Leu Met Thr Phe Ser Ile Ile Glu Leu Phe Ile Ser Leu Pro			
96	50 55 60			
97	Phe Ser Ile Leu Gly Cys His Ser Glu Asp Cys Asp Cys Glu Gln Cys			
98	65 70 75 80			
99	Cys			
102	<210> SEQ ID NO: 5			
103	<211> LENGTH: 417			
104	<212> TYPE: DNA			
105	<213> ORGANISM: Homo sapien			
107	<400> SEQUENCE: 5			
108	atggattcaa gcaccgcaca cagtccggtg tttctggat ttcctccaga aatcactgct		60	
109	tcagaatatg agtccacaga actttcagcc acgacctttt caactcaaag ccccttgcaa		120	
110	aaatttatttg ctagaaaaat gaaaatctta gggactatcc agatcctgtt tggattatgt		180	
111	accttttctt ttggagttat cttccctttc acctgttaa aaccatatcc aaggtttccc		240	
112	tttatatttc ttccaggata tccattctgg ggctctgtt tggattttttt ttctggagcc		300	
113	ttccataattt cagtggaaat gaaaaccaca gaaactctga tcaaaactac atttgggtt		360	
114	attctcacca aaatagtcag tgtaaggctg ttactgcct gttcttggga attttga		417	

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Input Set : A:\SEQLIST - LEX-0109-USA.txt
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116 <210> SEQ ID NO: 6
 117 <211> LENGTH: 138
 118 <212> TYPE: PRT
 119 <213> ORGANISM: Homo sapien
 121 <400> SEQUENCE: 6
 122 Met Asp Ser Ser Thr Ala His Ser Pro Val Phe Leu Val Phe Pro Pro
 123 1 5 10 15
 124 Glu Ile Thr Ala Ser Glu Tyr Glu Ser Thr Glu Leu Ser Ala Thr Thr
 125 20 25 30
 126 Phe Ser Thr Gln Ser Pro Leu Gln Lys Leu Phe Ala Arg Lys Met Lys
 127 35 40 45
 128 Ile Leu Gly Thr Ile Gln Ile Leu Phe Gly Ile Met Thr Phe Ser Phe
 129 50 55 60
 130 Gly Val Ile Phe Leu Phe Thr Leu Leu Lys Pro Tyr Pro Arg Phe Pro
 131 65 70 75 80
 132 Phe Ile Phe Leu Ser Gly Tyr Pro Phe Trp Gly Ser Val Leu Phe Ile
 133 85 90 95
 134 Asn Ser Gly Ala Phe Leu Ile Ala Val Lys Arg Lys Thr Thr Glu Thr
 135 100 105 110
 136 Leu Ile Lys Thr Thr Phe Val Val Ile Leu Thr Lys Ile Val Ser Val
 137 115 120 125
 138 Arg Leu Leu Leu Ser Cys Ser Trp Glu Phe
 139 130 135
 141 <210> SEQ ID NO: 7
 142 <211> LENGTH: 450
 143 <212> TYPE: DNA
 144 <213> ORGANISM: Homo sapien
 146 <400> SEQUENCE: 7
 147 atggattcaa gcaccgcaca cagtccggtg tttctggat ttcctccaga aatcactgct 60
 148 tcagaatatg agtccacaga actttcagcc acgacctttt caactcaaag ccccttgaa 120
 149 aaattatttg ctagaaaaat gaaaatctta gggactatcc agatcctgtt tggaattatg 180
 150 accttttctt ttggagttat cttcctttc accttggtaa aaccatatcc aaggtttccc 240
 151 ttatatttc ttccaggata tccattctgg ggctctgttt tgttcattaa ttctggagcc 300
 152 ttccataattg cagtggaaag aaaaaccaca gaaactctgg gaattttgat tacattgtg 360
 153 actttcagca ttattgaatt attcattttct ctgcctttct caattttggg gtgccactca 420
 154 gaggattgtt atgtgaaca atgttggta 450
 156 <210> SEQ ID NO: 8
 157 <211> LENGTH: 149
 158 <212> TYPE: PRT
 159 <213> ORGANISM: Homo sapien
 161 <400> SEQUENCE: 8
 162 Met Asp Ser Ser Thr Ala His Ser Pro Val Phe Leu Val Phe Pro Pro
 163 1 5 10 15
 164 Glu Ile Thr Ala Ser Glu Tyr Glu Ser Thr Glu Leu Ser Ala Thr Thr
 165 20 25 30
 166 Phe Ser Thr Gln Ser Pro Leu Gln Lys Leu Phe Ala Arg Lys Met Lys
 167 35 40 45
 168 Ile Leu Gly Thr Ile Gln Ile Leu Phe Gly Ile Met Thr Phe Ser Phe
 169 50 55 60

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170	Gly	Val	Ile	Phe	Leu	Phe	Thr	Leu	Leu	Lys	Pro	Tyr	Pro	Arg	Phe	Pro
																80
171	65				70					75						
172	Phe	Ile	Phe	Leu	Ser	Gly	Tyr	Pro	Phe	Trp	Gly	Ser	Val	Leu	Phe	Ile
																95
173					85					90						
174	Asn	Ser	Gly	Ala	Phe	Leu	Ile	Ala	Val	Lys	Arg	Lys	Thr	Thr	Glu	Thr
																110
175					100					105						
176	Leu	Gly	Ile	Leu	Ile	Thr	Leu	Met	Thr	Phe	Ser	Ile	Ile	Glu	Leu	Phe
																125
177					115					120						
178	Ile	Ser	Leu	Pro	Phe	Ser	Ile	Leu	Gly	Cys	His	Ser	Glu	Asp	Cys	Asp
179					130					135						140
180	Cys	Glu	Gln	Cys	Cys											
181	145															
183	<210>	SEQ	ID	NO:	9											
184	<211>	LENGTH:	676													
185	<212>	TYPE:	DNA													
186	<213>	ORGANISM:	Homo sapien													
188	<400>	SEQUENCE:	9													
189	tcatctcctt	tcaaatttac	accgacacca	tcatggattc	aagcaccgca	cacagtccgg										60
190	tgtttctggt	atttcctcca	gaaatcactg	cttcagaata	tgagtccaca	gaactttcag										120
191	ccacgacctt	ttcaactcaa	agccccttgc	aaaattatt	tgctagaaaa	ataaaaatct										180
192	tagggactat	ccagatcctg	tttggaaatta	tgacctttc	ttttggagtt	atttccttt										240
193	tcaccttgg	aaaaccatat	ccaaggttcc	cctttataatt	tctttcagga	tatccattct										300
194	ggggctctgt	tttggtcatt	aattctggag	ccttcctaat	tgcagtgaaa	agaaaaacca										360
195	cagaaactct	gataatattt	agccgaataa	tgaatcttct	tagtccctg	agagcaatag										420
196	ctggaatcat	tctcctcaca	tttggtttca	tcctagatca	aaactacatt	tgtggttatt										480
197	ctcacaaaaa	tagtcagtgt	aaggctgtta	ctgtcctgtt	cttgggaatt	ttgattacat										540
198	tcatgacttt	cagcattatt	gaattattca	tttctctgcc	tttctcaatt	ttggggtgcc										600
199	actcagagga	tttgttattgt	gaacaatgtt	gttgactagc	actgtgagaa	taaagatgtg										660
200	ttaaaataaa	aaaaaaa														676

VERIFICATION SUMMARY

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Input Set : A:\SEQLIST - LEX-0109-USA.txt

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L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date